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(2) Health and environmental effects data. All existing data regarding health and environmental effects of the microorganism must be reported in accordance with \$725.160.

§ 725.370 EPA review of the TME application.

General procedures for review of all submissions under this part are contained in §§ 725.28 through 725.60. In addition, the following procedures apply to EPA review of TME applications submitted under this subpart:

- (a) No later than 45 days after EPA receives a TME, the Agency will either approve or deny the application.
- (b) A submitter may only proceed with test marketing activities after receipt of EPA approval.
- (c) In approving a TME application, EPA may impose any restrictions necessary to ensure that the microorganism will not present an unreasonable risk of injury to health and the environment as a result of test marketing.

Subpart G—General Exemptions for New Microorganisms

§725.400 Scope and purpose.

- (a) This subpart describes exemptions from reporting under subpart D of this part, and from review under this part altogether, for manufacturing and importing of certain new microorganisms for commercial purposes.
- (b) Recipient microorganisms eligible for the tiered exemption from review under this part are listed in §725.420.
- (c) Criteria for the introduced genetic material contained in the new microorganisms are described in §725.421.
- (d) Physical containment and control technologies are described in §725.422.
- (e) The conditions for the Tier I exemption are listed in §725.424.
- (f) In lieu of complying with subpart D of this part, persons using recipient microorganisms eligible for the tiered exemption may submit a Tier II exemption request. The limited reporting requirements for the Tier II exemption, including data requirements, are described in §§ 725.450 and 725.455.
- (g) EPA review procedures for the Tier II exemption are set forth in \$725.470.

(h) Subparts A through C of this part apply to any submission under this subpart.

§ 725.420 Recipient microorganisms.

The following recipient microorganisms are eligible for either exemption under this subpart:

- (a) Acetobacter aceti.
- $\hbox{(b) } \textit{Aspergillus niger}.$
- (c) Aspergillus oryzae.
- ${\rm (d)}\ Bacillus\ licheniform is.$
- (e) Bacillus subtilis.
- (f) Clostridium acetobutylicum.
- (g) Escherichia coli K-12.
- $(h) \ Penicillium \ roque forti.$
- (i) Saccharomyces cerevisiae.
- (j) Saccharomyces uvarum.

§ 725.421 Introduced genetic material.

For a new microorganism to qualify for either exemption under this subpart, introduced genetic material must meet all of the criteria listed in this section.

- (a) *Limited in size*. The introduced genetic material must consist only of the following:
 - (1) The structural gene(s) of interest.
- (2) The regulatory sequences permitting the expression of solely the gene(s) of interest.
- (3) Associated nucleotide sequences needed to move genetic material, including linkers, homopolymers, adaptors, transposons, insertion sequences, and restriction enzyme sites.
- (4) The nucleotide sequences needed for vector transfer.
- (5) The nucleotide sequences needed for vector maintenance.
- (b) Well-characterized. For introduced genetic material, well-characterized means that the following have been determined:
- (1) The function of all of the products expressed from the structural gene(s).
- (2) The function of sequences that participate in the regulation of expression of the structural gene(s).
- (3) The presence or absence of associated nucleotide sequences and their associated functions, where associated nucleotide sequences are those sequences needed to move genetic material including linkers, homopolymers, adaptors, transposons, insertion sequences, and restriction enzyme sites.

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- (c) Poorly mobilizable. The ability of the introduced genetic material to be transferred and mobilized is inactivated, with a resulting frequency of transfer of less than 10^{-8} transfer events per recipient.
- (d) Free of certain sequences. (1) The introduced genetic material must not contain a functional portion of any of the toxin-encoding sequences described in this paragraph (d).
- (i) For the purposes of this section, a functional portion of a toxin-encoding sequence means any sequence which codes for a polypeptide that has one of the following effects:
- (A) It directly or indirectly contributes to toxic effects in humans. Directly contributes to toxic effects in humans means those sequences encoding polypeptides that have direct toxicity to target cells. An example of a sequence which directly contributes to toxic effects in humans is one which encodes the portion of diphtheria toxin, listed in paragraph (d)(2) of this section, capable of interacting with elongation factor 2, leading to inhibition of protein synthesis in target respiratory, heart, kidney, and nerve tissues. Indirectly contributes to toxic effects in humans means a sequence whose encoded polypeptide is not directly toxic to target cells, yet still adversely affects humans. An example of a sequence which indirectly contributes to toxic effects is the sequence which encodes the portion of the botulinum toxin, listed in paragraph (d)(3) of this section, capable of blocking the of acetylcholine gangliosides. Botulinum toxin affects neuromuscular junctions by its blockage of acetylcholine release, leading to irreversible relaxation of muscles and respiratory arrest.
- (B) It binds a toxin or toxin precursor to target human cells.
- (C) It facilitates intracellular transport of a toxin in target human cells.
- (ii) While these toxins are listed (with synonyms in parentheses) in paragraphs (d)(2) through (d)(7) of this section according to the source organism, it is use of the nucleotide sequences that encode the toxins that is being restricted and not the use of the source organisms. The source organisms are listed to provide specificity

in identification of sequences whose use is restricted. Although similar or identical sequences may be isolated from organisms other than those listed below in paragraphs (d)(2) through (d)(7) of this section, these comparable toxin sequences, regardless of the organism from which they are derived, must not be included in the introduced genetic material.

(2) Sequences for protein synthesis inhibitor.

Sequence Source

Toxin Name

Corvnebacterium diphtheriae & C. ulcerans Pseudomonas aeruginosa Shigella dysenteriae

Exotoxin A Shigella toxin (Shiga toxin, Shigella dysenteriae type I toxin, Vero cell toxin)

Diphtheria toxin

Abrin

Abrus precatorius, seeds Ricinus communis, seeds

(3) Sequences for neurotoxins.

Sequence Source

Toxin Name

Clostridium botulinum Clostridium tetani

Proteus mirabilis Staphylococcus aureus Yersinia pestis

Snake toxins Bungarus caeruleus Bungarus multicinctus

Crotalus spp. Dendroaspis viridis Naia naia varieties Notechia scutatus Oxyuranus scutellatus Invertebrate toxins

Chironex fleckeri Androctnus australis Centruroides sculpturatus

(4) Sequences cytolysins.

Neurotoxins A, B, C1, D, E, F. G (Botulinum toxins. botulinal toxins) Tetanus toxin (tetanospasmin) Neurotoxin Alpha toxin (alpha lysin) Murine toxin

Caeruleotoxin Beta-bungarotoxin (phospholipase) Crotoxin (phospholipase) Neurotoxin Neurotoxin Notexin (phospholipase) Taipoxin

Neurotoxin Neurotoxin Neurotoxin

Alveolysin

Cereolysin

Laterosporolysin

oxuaen for

Sequence Source

Toxin Name

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Bacillus alve Bacillus cereus Bacillus laterosporus Bacillus thuringiensis Clostridium bifermentans Clostridium botulinum Clostridium caproicum Clostridium chauvoei Clostridium histolyticum Clostridium novyi Clostridium oedematiens Clostridium perfringens Clostridium senticum Clostridium sordellii Clostridium tetani Listeria monocytogenes Streptococcus pneumoniae Streptococcus pyogene

Thuringiolysin Lysin Lysin Delta-toxin Epsilon-toxin . Gamma-toxin Delta-toxin Theta-toxin (Perfringolysin) Delta-toxin Lysin **Tetanolysin** Listeriolysin (A B) Pneumolysin Streptolysin O (SLO)

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(5) Sequences for toxins affecting membrane function.

Sequence Source

Toxin Name

Bacillus anthracis

Bacillus cereus

Bordetella pertussis

Clostridium botulinum Clostridium difficile Clostridium perfringens Escherichia coli & other Enterobacteriaceae spp

Legionella pneumophila Vibrio cholerae & Vibrio mimicus

Edema factor (Factors I II): Lethal factor (Factors II III) Enterotoxin (diarrheagenic toxin, mouse lethal factor)

Adenylate cyclase (Heat-labile factor): Pertussigen (pertussis toxin, islet activating factor, histamine sensitizing factor. lymphocytosis promoting

C2 toxin Enterotoxin (toxin A) Beta-toxin; Delta-toxin Heat-labile enterotoxins (LT); Heat-stable enterotoxins (STa, ST1 subtypes ST1a ST1b; also STb, STII)

Cholera toxin (choleragen)

(6) Sequences that affect membrane integrity.

Sequence Source

Toxin Name

Clostridium hifermentans & other Clostridium spp Clostridium perfringens

Corynebacterium pyogenes &

other Corynebacterium spp.

Staphylococcus aureus

(7)Sequences cytotoxins.

Lecithinase

Alpha-toxin (phospholipase C. lecithinase): Enterotoxin Cytolysin (phospholipase C), Ovis toxin (sphingomyelinase D)

Beta-lysin (beta toxin)

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Sequence Source

Adenia digitata Aeromonas hydrophila

Clostridium difficile Clostridium perfringens

Escherichia coli & other Enterobacteriaceae spp. Pseudomonas aeruginosa Staphylococcus aureus

Staphylococcus aureus & Pseudomonas aeruginosa Streptococcus pyogenes

Toxin Name

Modeccin Aerolysin (beta-lysin, cytotoxic lysin) Cytotoxin (toxin B) Beta-toxin; Epsilon-toxin; Kappa-toxin Cytotoxin (Shiga-like toxin, Vero cell toxin) Proteases Gamma lysin (Gamma toxin); Enterotoxins (SEA, SEB, SEC. SED SEE); Pyrogenic exotoxins A B: Toxic shock syndrome toxins (TSST-1) Leucocidin (leukocidin,

cytotoxin) Streptolysin S (SLS); Erythrogenic toxins (scarlet fever toxins, pyrogenic exotoxins) Heat-stable enterotoxins (ST)

Yersinia enterocolitica

§725.422 Physical containment and control technologies.

The manufacturer must meet all of the following criteria for physical containment and control technologies for any facility in which the new microorganism will be used for a Tier I exemption; these criteria also serve as guidance for a Tier II exemption.

- (a) Use a structure that is designed and operated to contain the new microorganism.
 - (b) Control access to the structure.
- (c) Provide written, published, and implemented procedures for the safety of personnel and control of hygiene.
- (d) Use inactivation procedures demonstrated and documented to be effective against the new microorganism contained in liquid and solid wastes prior to disposal of the wastes. The inactivation procedures must reduce viable microbial populations by at least 6 logs in liquid and solid wastes.
- (e) Use features known to be effective in minimizing viable microbial populations in aerosols and exhaust gases released from the structure, and document use of such features.
- (f) Use systems for controlling dissemination of the new microorganism through other routes, and document use of such features.
- (g) Have in place emergency clean-up procedures.

§725.424 Requirements for the Tier I exemption.

- (a) Conditions of exemption. The manufacture or import of a new microorganism for commercial purposes is not subject to review under this part if all of the following conditions are met for all activities involving the new microorganism:
- (1) The recipient microorganism is listed in and meets any requirements specified in §725.420.
- (2) The introduced genetic material meets the criteria under §725.421.
- (3) The physical containment and control technologies of any facility in which the microorganism will be manufactured, processed, or used meet the criteria under §725.422.
- (4) The manufacturer or importer submits a certification described in paragraph (b) of this section to EPA at least 10 days before commencing initial manufacture or import of a new microorganism derived from a recipient microorganism listed in §725.420.